## Figure 1

Ser Trp Phe Pro Val Gln Gly Pro Ala Asp Ile Cys Gln Cys Cys Asn Lys Gly Asp Gly Thr 308 318 328

Pro Ser His Ser Arg Arg Glu Pro His Val Met Ser Gln Trp Ser Arg Ser Val Ser

338

1

348

Table 1

Position No. Preferred Amino Acid		Other Representative Amino Acids		
310 315 320 323 324 326 328 329 330 332 334 335 337 339 341 342 345	phe pro gln asn lys asp gly thr pro his arg arg pro val ser gln arg	tyr  asp gln  asn, glu, gln, ile, pro, phe, cys ala, ile ser  lys lys lys met iso, met thr asn lys		
	_	•		

Figure 2

Name: HUMAN Length: 41 SWFPVQGPADICQCCNKGDCGTPSHSRRQPHVMSQWSRSVS

Name HUMAN	Residue Thr	No. 22	Potential 0.0285	Threshold 0.4982	Assignment .
Name	Residue	No	Potential		
HUMAN				Threshold	Assignment
	Ser	1	0.0629	0.5132	_
HUMAN	Ser	24	0.0104	0.5253	
HUMAN	Ser	26			
		26	0.0265	0.5309	
HUMAN	Ser	34	0.0033	0.6267	
HUMAN	Ser	37	- <del>-</del>		
HUMAN			0.4498	0.5825	
	Ser	39	0.0009	0.5126	
HUMAN	Ser	41	0.0082		
			0.0082	0.5022	•

Figure 3

Name: MOUSE Length: 41

SWLPVQGDADICDCCSHGNCSNSSSSQFQIHGPRQWSKLVS

Name	Residue	No.	Potential	Threshold	Assignment
Name	Residue	No.	Potential	Threshold	Assignment
MOUSE	Ser	1	0.0116	0.5078	
MOUSE	Ser	16	0.0002	0.5747	
MOUSE	Ser	21	0.0403	0.5845	
MOUSE	Ser	23	0.0074	0.5310	
MOUSE	Ser	24	0.1625	0.5397	
MOUSE	Ser	25	0.0167	0.5585	
MOUSE	Ser	26	0.0003	0.5656	
MOUSE	Ser	37	0.0024	0.5627	
MOUSE	Ser	41	0.0005	0.5127	